

*Sub E1)*

*D1  
cont* 2. (Twice Amended) A recombinant host cell comprising a nucleic acid molecule comprising the nucleotide sequence of any of Claims 1, 39, 40, or 47.

*D2 Sub E2)* 3. (Amended) A vector comprising the nucleic acid molecule of Claims 1, 39, 40, or 47.

*D3* 13. (Twice Amended) A process for producing an FGF-like polypeptide comprising culturing the host cell of Claim 9 under suitable conditions to express the polypeptide, wherein said polypeptide can be isolated from the culture.

39. (Twice Amended) An isolated nucleic acid molecule comprising:

- (a) a region of the nucleotide sequence of SEQ ID NO: 3, encoding a polypeptide fragment of at least about 25 amino acid residues, wherein the polypeptide fragment has an activity of the polypeptide set forth in SEQ ID NO: 4, or is antigenic;
- (b) a region of the nucleotide sequence of SEQ ID NO: 3 comprising a fragment of at least about 16 nucleotides; or
- (c) a nucleotide sequence complementary to the nucleotide sequence of either (a) or (b).

40. (Twice Amended) An isolated nucleic acid molecule comprising:

- (a) a nucleotide sequence encoding a polypeptide as set forth in SEQ ID NO: 4 with at least one conservative amino acid substitution, wherein the encoded polypeptide has an activity of the polypeptide set forth in SEQ ID NO: 4;
- (b) a region of the nucleotide sequence of (a) comprising a fragment of at least about 16 nucleotides; or
- (c) a nucleotide sequence complementary to the nucleotide sequence of either (a) or (b).

42. (Amended) A process of producing an FGF-like polypeptide comprising culturing the recombinant host cell of Claim 9 under suitable conditions to express the polypeptide.

Please add the following new claim.

47. An isolated nucleic acid molecule comprising:

(a) a nucleotide sequence encoding a polypeptide comprising the amino acid sequence:

Met Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Gly Leu Trp Val Xaa Xaa Xaa  
Leu Ala Xaa Xaa Leu Leu Gly Xaa Xaa Gln Ala Xaa Pro Ile Pro Asp Ser Ser Pro  
Leu Leu Gln Phe Gly Gly Gln Val Arg Gln Arg Tyr Leu Tyr Thr Asp Asp Xaa  
Gln Xaa Thr Glu Ala His Leu Glu Ile Arg Glu Asp Gly Thr Val Xaa Gly Ala Ala  
Xaa Xaa Ser Pro Glu Ser Leu Leu Xaa Leu Lys Ala Leu Lys Pro Gly Val Ile Gln  
Ile Leu Gly Val Lys Xaa Ser Arg Phe Leu Cys Gln Xaa Pro Asp Gly Ala Leu Tyr  
Gly Ser Xaa His Phe Asp Pro Glu Ala Cys Ser Phe Arg Glu Leu Leu Leu Asp  
Gly Tyr Asn Val Tyr Gln Ser Glu Ala His Gly Leu Pro Leu Xaa Leu Pro Xaa Xaa  
Xaa Ser Pro Xaa Xaa Asp Xaa Xaa Xaa Xaa Gly Pro Xaa Arg Phe Leu Pro Xaa  
Pro Gly Leu Xaa Xaa Xaa Pro Xaa Xaa Xaa Xaa Gly Xaa Leu Xaa Pro Xaa Pro  
Pro Asp Val Gly Ser Ser Asp Pro Leu Ser Met Val Xaa Pro Xaa Gln Gly Arg Ser  
Pro Ser Tyr Ala Ser,

wherein the residue at either position 2 or 177 may be either aspartic acid or glutamic acid;

the residue at position 6 may be either threonine or serine;

the residue at position 18 may be either leucine or valine;

the residue at any of positions 76, 105, or 155 may be either arginine or glutamine;

the residue at either position 83 or 185 may be either glutamic acid or glutamine;

the residue at either position 99 or 158 may be either threonine or alanine;

the residue at position 146 may be either histidine or arginine;

the residue at position 154 may be either histidine or asparagine;

the residue at position 168 may be either leucine or methionine;